SEQUENCE LISTING

<110> John P. Carulli et al.

<130> 032796-014

<120> THE HIGH BONE MASS GENE OF 11q13.3

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Met Glu Ala Ala Pro Pro Gly Pro Pro Trp Pro Leu Leu	
1 5 10	
ctg ctg ctg ctg ctg ctg gcg ctg tgc ggc tgc ccg gcc ccc gcc	157
Leu Leu Leu Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala	
15 20 25	
gcg gcc tcg ccg ctc ctg cta ttt gcc aac cgc cgg gac gta cgg ctg	205
Ala Ala Ser Pro Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu	
30 35 40 45	

The first production of the first of the control of

190					195					200					205	
gag	cag	aag	ctc	tac	tgg	gct	gac	gcc	aag	ctc	agc	ttc	atc	cac	cgt	733
Glu	Gln	Lys	Leu	Tyr	Trp	Ala	Asp	Ala	Lys	Leu	Ser	Phe	Ile	His	Arg	
				210					215					220		
gcc	aac	ctg	gac	ggc	tcg	ttc	cgg	cag	aag	gtg	gtg	gag	ggc	agc	ctg	781
Ala	Asn	Leu	Asp	Gly	Ser	Phe	Arg	Gln	Lys	Val	Val	Glu	Gly	Ser	Leu	
			225					230					235			
acg	cac	ccc	ttc	gcc	ctg	acg	ctc	tcc	999	gac	act	ctg	tac	tgg	aca	829
Thr	His	Pro	Phe	Ala	Leu	Thr	Leu	Ser	Gly	Asp	Thr	Leu	Tyr	Trp	Thr	
		240					245					250				
gac	tgg	cag	acc	cgc	tcc	atc	cat	gcc	tgc	aac	aag	cgc	act	aaa	ggg	877
Asp	Trp	Gln	Thr	Arg	Ser	Ile	His	Ala	Cys	Asn	Lys	Arg	Thr	Gly	Gly	
	255					260					265					
aag	agg	aag	gag	atc	ctg	agt	gcc	ctc	tac	tca	ccc	atg	gac	atc	cag	925
Lys	Arg	Lys	Glu	Ile	Leu	Ser	Ala	Leu	Tyr	Ser	Pro	Met	Asp	Ile	Gln	
270					275					280					285	
gtg	ctg	agc	cag	gag	cgg	cag	cct	ttc	ttc	cac	act	cgc	tgt	gag	gag	973
Val	Leu	Ser	Gln	Glu	Arg	Gln	Pro	Phe	Phe	His	Thr	Arg	Cys	Glu	Glu	
				290					295					300		
gac	aat	ggc	ggc	tgc	tcc	cac	ctg	tgc	ctg	ctg	tcc	cca	agc	gag	cct	1021
Asp	Asn	Gly	Gly	Cys	Ser	His	Leu	Cys	Leu	Leu	Ser	Pro	Ser	Glu	Pro	
			305					310					315			
									gtg							1069
Phe	Tyr		Cys	Ala	Cys	Pro		Gly	Val	Gln	Leu	Gln	Asp	Asn	Gly	
		320					325					330				
									gtg							1117
Arg		Сув	Lys	Ala	Gly		Glu	Glu	Val	Leu		Leu	Ala	Arg	Arg	
	335					340					345					
acg	gac	cta	cgg	agg	atc	tcg	ctg	gac	acg	ccg	gac	ttc	acc	gac	atc	1165

Thi	Ası	p Lev	ı Arç	g Arg	j Ile	e Ser	Leu	Asp	Thr	Pro	Asp	Phe	e Thi	Asp	lle	
350					355	5				360)				365	
gto	g ctg	g cag	g gto	g gad	gac	ato	: cgg	cac	gcc	att	geo	ato	gac	tac	gac	1213
Val	. Lev	ı Glr	ı Val	Asp	Asp	Ile	Arg	His	Ala	Ile	Ala	ıle	. Asp	туг	Asp	
				370)				375					380)	
ccg	r cta	gag	ggc	tat	gto	tac	tgg	aca	gat	gac	gag	gtg	cgg	gcc	atc	1261
Pro	Leu	Glu	Gly	Tyr	Val	Tyr	Trp	Thr	Asp	Asp	Glu	Val	Arg	Ala	Ile	
			385					390					395			
cgc	agg	gcg	tac	ctg	gac	aaa	tct	999	gcg	cag	acg	ctg	gtc	aac	acc	1309
Arg	Arg	Ala	Tyr	Leu	Asp	Gly	Ser	Gly	Ala	Gln	Thr	Leu	Val	Asn	Thr	
		400					405					410				
gag	atc	aac	gac	ccc	gat	ggc	atc	gcg	gtc	gac	tgg	gtg	gcc	cga	aac	1357
Glu	Ile	Asn	Asp	Pro	Asp	Gly	Ile	Ala	Val	Asp	Trp	Val	Ala	Arg	Asn	
	415					420					425					
ctc	tac	tgg	acc	gac	acg	ggc	acg	gac	cgc	atc	gag	gtg	acg	cgc	ctc	1405
Leu	Tyr	Trp	Thr	Asp	Thr	Gly	Thr	Asp	Arg	Ile	Glu	Val	Thr	Arg	Leu	
430					435					440					445	
aac	ggc	acc	tcc	cgc	aag	atc	ctg	gtg	tcg	gag	gac	ctg	gac	gag	ccc	1453
Asn	Gly	Thr	Ser	Arg	Lys	Ile	Leu	Val	Ser	Glu	Asp	Leu	Asp	Glu	Pro	
				450					455					460		
cga	gcc	atc	gca	ctg	cac	ccc	gtg	atg	ggc	ctc	atg	tac	tgg	aca	gac	1501
Arg	Ala	Ile	Ala	Leu	His	Pro	Val	Met	Gly	Leu	Met	Tyr	Trp	Thr	Asp	
			465					470					475			
tgg	gga	gag	aac	cct	aaa	atc	gag	tgt	gcc	aac	ttg	gat	ggg	cag	gag	1549
Trp	Gly	Glu	Asn	Pro	Lys	Ile	Glu	Cys	Ala	Asn	Leu	Asp	Gly	Gln	Glu	
		480					485					490				
cgg	cgt	gtg	ctg	gtc	aat	gcc	tcc	ctc	a aa	tgg	ccc	aac	ggc	ctg	gcc	1597
Arg	Arg	Val	Leu	Val	Asn	Ala	Ser	Leu	Gly	Trp	Pro	Asn	Gly	Leu	Ala	
	495					500					505					

ctg	gac	ctg	cag	gag	9 99	aag	ctc	tac	tgg	gga	gac	gcc	aag	aca	gac	16	545
Leu	Asp	Leu	Gln	Glu	Gly	Lys	Leu	Tyr	Trp	Gly	Asp	Ala	Lys	Thr	Asp		
510					515					520					525		
aag	atc	gag	gtg	atc	aat	gtt	gat	999	acg	aag	agg	cgg	acc	ctc	ctg	16	593
Lys	Ile	Glu	Val	Ile	Asn	Val	Asp	Gly	Thr	Lys	Arg	Arg	Thr	Leu	Leu		
				530					535					540			
gag	gac	aag	ctc	ccg	cac	att	ttc	999	ttc	acg	ctg	ctg	999	gac	ttc	17	741
Glu	Asp	Lys	Leu	Pro	His	Ile	Phe	Gly	Phe	Thr	Leu	Leu	Gly	Asp	Phe		
			545					550					555				
atc	tac	tgg	act	gac	tgg	cag	cgc	cgc	agc	atc	gag	cgg	gtg	cac	aag	17	789
Ile	Tyr	Trp	Thr	Asp	Trp	Gln	Arg	Arg	Ser	Ile	Glu	Arg	Val	His	Lys		
		560					565					570					
gtc	aag	gcc	agc	cgg	gac	gtc	atc	att	gac	cag	ctg	ccc	gac	ctg	atg	18	337
Val	Lys	Ala	Ser	Arg	Asp	Val	Ile	Ile	Asp	Gln	Leu	Pro	Asp	Leu	Met		
	575					580					585						
999	ctc	aaa	gct	gtg	aat	gtg	gcc	aag	gtc	gtc	gga	acc	aac	ccg	tgt	18	385
Gly	Leu	Lys	Ala	Val	Asn	Val	Ala	Lys	Val	Val	Gly	Thr	Asn	Pro	Cys		
590					595					600					605		
gcg	gac	agg	aac	999	999	tgc	agc	cac	ctg	tgc	ttc	ttc	aca	ccc	cac	19	933
Ala	Asp	Arg	Asn	Gly	Gly	Cys	Ser	His	Leu	Cys	Phe	Phe	Thr	Pro	His		
				610					615					620			
gca	acc	cgg	tgt	ggc	tgc	ccc	atc	ggc	ctg	gag	ctg	ctg	agt	gac	atg	19	981
Ala	Thr	Arg	Cys	Gly	Cys	Pro	Ile	Gly	Leu	Glu	Leu	Leu	Ser	Asp	Met		
			625					630					635				
aag	acc	tgc	atc	gtg	cct	gag	gcc	ttc	ttg	gtc	ttc	acc	agc	aga	gcc	20	29
Lys	Thr	Cys	Ile	Val	Pro	Glu	Ala	Phe	Leu	Val	Phe	Thr	Ser	Arg	Ala		
		640					645					650					
gcc	atc	cac	agg	atc	tcc	ctc	gag	acc	aat	aac	aac	gac	gtg	gcc	atc	20	77
Ala	Ile	His	Arg	Ile	Ser	Leu	Glu	Thr	Asn	Asn	Asn	Asp	Val	Ala	Ile		

ccg ctc acg ggc gtc aag gag gcc tca gcc ctg gac ttt gat gtg tcc Pro Leu Thr Gly Val Lys Glu Ala Ser Ala Leu Asp Phe Asp Val Ser aac aac cac atc tac tgg aca gac gtc agc ctg aag acc atc agc cgc Asn Asn His Ile Tyr Trp Thr Asp Val Ser Leu Lys Thr Ile Ser Arg gcc ttc atg aac ggg agc tcg gtg gag cac gtg gtg gag ttt ggc ctt Ala Phe Met Asn Gly Ser Ser Val Glu His Val Val Glu Phe Gly Leu gac tac ccc gag ggc atg gcc gtt gac tgg atg ggc aag aac ctc tac Asp Tyr Pro Glu Gly Met Ala Val Asp Trp Met Gly Lys Asn Leu Tyr tgg gcc gac act ggg acc aac aga atc gaa gtg gcg cgg ctg gac ggg Trp Ala Asp Thr Gly Thr Asn Arg Ile Glu Val Ala Arg Leu Asp Gly cag ttc cgg caa gtc ctc gtg tgg agg gac ttg gac aac ccg agg tcg Gln Phe Arg Gln Val Leu Val Trp Arg Asp Leu Asp Asn Pro Arg Ser ctg gcc ctg gat ccc acc aag ggc tac atc tac tgg acc gag tgg ggc Leu Ala Leu Asp Pro Thr Lys Gly Tyr Ile Tyr Trp Thr Glu Trp Gly ggc aag ccg agg atc gtg cgg gcc ttc atg gac ggg acc aac tgc atg Gly Lys Pro Arg Ile Val Arg Ala Phe Met Asp Gly Thr Asn Cys Met acg ctg gtg gac aag gtg ggc cgg gcc aac gac ctc acc att gac tac Thr Leu Val Asp Lys Val Gly Arg Ala Asn Asp Leu Thr Ile Asp Tyr get gae cag ege ete tae tgg ace gae etg gae ace aac atg ate gag

Ala	Asp	Gln	Arg	Leu	Tyr	Trp	Thr	Asp	Leu	Asp	Thr	Asn	Met	Ile	Glu	
	815					820					825					
tcg	tcc	aac	atg	ctg	ggt	cag	gag	cgg	gtc	gtg	att	gcc	gac	gat	ctc	2605
Ser	Ser	Asn	Met	Leu	Gly	Gln	Glu	Arg	Val	Val	Ile	Ala	Asp	Asp	Leu	
830					835					840					845	
ccg	cac	ccg	ttc	ggt	ctg	acg	cag	tac	agc	gat	tat	atc	tac	tgg	aca	2653
Pro	His	Pro	Phe	Gly	Leu	Thr	Gln	Tyr	Ser	Asp	Tyr	Ile	Tyr	Trp	Thr	
				850					855					860		
gac	tgg	aat	ctg	cac	agc	att	gag	cgg	gcc	gac	aag	act	agc	ggc	cgg	2701
Asp	Trp	Asn	Leu	His	Ser	Ile	Glu	Arg	Ala	Asp	Lys	Thr	Ser	Gly	Arg	
			865					870					875			
aac	cgc	acc	ctc	atc	cag	ggc	cac	ctg	gac	ttc	gtg	atg	gac	atc	ctg	2749
Asn	Arg	Thr	Leu	Ile	Gln	Gly	His	Leu	Asp	Phe	Val	Met	Asp	Ile	Leu	
		880					885					890				
gtg	ttc	cac	tcc	tcc	cgc	cag	gat	ggc	ctc	aat	gac	tgt	atg	cac	aac	2797
Val	Phe	His	Ser	Ser	Arg	Gln	Asp	Gly	Leu	Asn	Asp	Cys	Met	His	Asn	
	895					900					905					
aac	999	cag	tgt	999	cag	ctg	tgc	ctt	gcc	atc	ccc	ggc	ggc	cac	cgc	2845
Asn	Gly	Gln	Cys	Gly	Gln	Leu	Cys	Leu	Ala	Ile	Pro	Gly	Gly	His	Arg	
910					915					920					925	
tgc	ggc	tgc	gcc	tca	cac	tac	acc	ctg	gac	ccc	agc	agc	cgc	aac	tgc	2893
Cys	Gly	Cys	Ala	Ser	His	Tyr	Thr	Leu	Asp	Pro	Ser	Ser	Arg	Asn	Cys	
				930					935					940		
agc	ccg	ccc	acc	acc	ttc	ttg	ctg	ttc	agc	cag	aaa	tct	gcc	atc	agt	2941
Ser	Pro	Pro	Thr	Thr	Phe	Leu	Leu	Phe	Ser	Gln	Lys	Ser	Ala	Ile	Ser	
			945					950					955			
cgg	atg	atc	ccg	gac	gac	cag	cac	agc	ccg	gat	ctc	atc	ctg	ccc	ctg	2989
Arg	Met	Ile	Pro	Asp	Asp	Gln	His	Ser	Pro	Asp	Leu	Ile	Leu	Pro	Leu	
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ca	t gga	ctg	agg	aac	gtc	aaa	gcc	atc	gac	tat	gac	сса	ctg	gac	aag	3037
Hi	s Gly	Leu	Arg	Asn	Val	Lys	Ala	Ile	Asp	Tyr	Asp	Pro	Leu	Asp	Lys	
	975					980					985					
tt	c atc	tac	tgg	gtg	gat	999	cgc	cag	aac	atc	aag	cga	gcc	aag	gac	3085
Ph	e Ile	Tyr	Trp	Val	Asp	Gly	Arg	Gln	Asn	Ile	Lys	Arg	Ala	Lys	Asp	
99	0				995					1000)				1005	
ga	c ggg	acc	cag	ccc	ttt	gtt	ttg	acc	tct	ctg	agc	caa	ggc	caa	aac	3133
As	o Gly	Thr	Gln	Pro	Phe	Val	Leu	Thr	Ser	Leu	Ser	Gln	Gly	Gln	Asn	
				1010)				1015	5				1020)	
CC	a gac	agg	cag	ccc	cac	gac	ctc	agc	atc	gac	atc	tac	agc	cgg	aca	3181
Pr	o Asp	Arg	Gln	Pro	His	Asp	Leu	Ser	Ile	Asp	Ile	Tyr	Ser	Arg	Thr	
			1025	5				1030)				1035	5		
ct	g ttc	tgg	acg	tgc	gag	gcc	acc	aat	acc	atc	aac	gtc	cac	agg	ctg	3229
Le	ı Phe	Trp	Thr	Cys	Glu	Ala	Thr	Asn	Thr	Ile	Asn	Val	His	Arg	Leu	
		1040)				1045	5				1050)			
ag	ggg	gaa	gcc	atg	ggg	gtg	gtg	ctg	cgt	999	gac	cgc	gac	aag	ccc	3277
Se	r Gly	Glu	Ala	Met	Gly	Val	Val	Leu	Arg	Gly	Asp	Arg	Asp	Lys	Pro	
	105	5				1060)				1065	5				
ag	g gcc	atc	gtc	gtc	aac	gcg	gag	cga	9 99	tac	ctg	tac	ttc	acc	aac	3325
Ar	g Ala	Ile	Val	Val	Asn	Ala	Glu	Arg	Gly	Tyr	Leu	Tyr	Phe	Thr	Asn	
10	70				1075	5				1080)				1085	
at	g cag	gac	cgg	gca	gcc	aag	atc	gaa	cgc	gca	gcc	ctg	gac	ggc	acc	3373
Me	Gln	Asp	Arg	Ala	Ala	Lys	Ile	Glu	Arg	Ala	Ala	Leu	Asp	Gly	Thr	
				1090)				1095	5				1100)	
ga	g cgc	gag	gtc	ctc	ttc	acc	acc	ggc	ctc	atc	cgc	cct	gtg	gcc	ctg	3421
Gl	_	C111	7/-1	LAII	Phe	Thr	Thr	Gly	Leu	Ile	Arg	Pro	Val	Ala	Leu	
	ı Arg	Giu														
			1105	5				1110					1115			
gt	a Arg	gac	1105 aac	aca	ctg	ggc		ctg	ttc				gcg	gac		3469

1120		1125		1130	
aag cgc att gag	agc tgt gac	ctg tca	ggg gcc aac	c cgc ctg acc	ctg 3517
Lys Arg Ile Glu	Ser Cys Asp	Leu Ser	Gly Ala Asn	n Arg Leu Thr	Leu
1135	114	0	114	45	
gag gac gcc aac	atc gtg cag	cct ctg	ggc ctg acc	c atc ctt ggc	aag 3565
Glu Asp Ala Asr	Ile Val Gln	Pro Leu	Gly Leu Thr	r Ile Leu Gly	Lys
1150	1155		1160		1165
cat ctc tac tgg	atc gac cgc	cag cag	cag atg atc	c gag cgt gtg	gag 3613
His Leu Tyr Trp	lle Asp Arg	Gln Gln	Gln Met Ile	e Glu Arg Val	Glu
	1170		1175	1180)
aag acc acc ggg	gac aag cgg	act cgc	atc cag ggc	c cgt gtc gcc	cac 3661
Lys Thr Thr Gly	Asp Lys Arg	Thr Arg	Ile Gln Gly	y Arg Val Ala	His
118	5	1190		1195	
ctc act ggc atc	cat gca gtg	gag gaa g	gtc agc ctg	g gag gag ttc	tca 3709
Leu Thr Gly Ile	His Ala Val	Glu Glu '	Val Ser Leu	ı Glu Glu Phe	Ser
1200		1205		1210	
gcc cac cca tgt	gcc cgt gac	aat ggt g	ggc tgc tcc	c cac atc tgt	att 3757
Ala His Pro Cys	Ala Arg Asp	Asn Gly (Gly Cys Ser	r His Ile Cys	Ile
1215	122	0	122	25	
gcc aag ggt gat	ggg aca cca	cgg tgc 1	tca tgc cca	a gtc cac ctc	gtg 3805
Ala Lys Gly Asp	Gly Thr Pro	Arg Cys S	Ser Cys Pro	o Val His Leu	Val
1230	1235		1240		1245
ctc ctg cag aac	-				
Leu Leu Gln Asn					
	1250		1255	1260	
gac cag ttt gca					
Asp Gln Phe Ala	-		lle Asp Cys		Ala
126	5	1270		1275	
tgg cgc tgt gad					gag 3949

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Trp 2	Arg	Cys	Asp	Gly	Phe	Pro	Glu	Cys	Asp	Asp	Gln	Ser	Asp	Glu	Glu	
		1280)				128	5				1290)			
ggc	tgc	ccc	gtg	tgc	tcc	gcc	gcc	cag	ttc	ccc	tgc	gcg	cgg	ggt	cag	3997
Gly	Cys	Pro	Val	Cys	Ser	Ala	Ala	Gln	Phe	Pro	Cys	Ala	Arg	Gly	Gln	
:	1295	5				1300)				1305	5				
tgt (gtg	gac	ctg	cgc	ctg	cgc	tgc	gac	ggc	gag	gca	gac	tgt	cag	gac	4045
Cys '	Val	Asp	Leu	Arg	Leu	Arg	Cys	Asp	Gly	Glu	Ala	Asp	Cys	Gln	Asp	
1310					1315	5				1320)				1325	
cgc 1	tca	gac	gag	gtg	gac	tgt	gac	gcc	atc	tgc	ctg	ccc	aac	cag	ttc	4093
Arg s	Ser	Asp	Glu	Val	Asp	Cys	Asp	Ala	Ile	Cys	Leu	Pro	Asn	Gln	Phe	
				1330)				1335	5				1340)	
cgg 1	tgt	gcg	agc	ggc	cag	tgt	gtc	ctc	atc	aaa	cag	cag	tgc	gac	tcc	4141
Arg (Cys	Ala	Ser	Gly	Gln	Cys	Val	Leu	Ile	Lys	Gln	Gln	Cys	Asp	Ser	
			1345	5				1350)				1355	5		
ttc	ccc	gac	tgt	atc	gac	ggc	tcc	gac	gag	ctc	atg	tgt	gaa	atc	acc	4189
Phe 1	Pro	Asp	Cys	Ile	Asp	Gly	Ser	Asp	Glu	Leu	Met	Cys	Glu	Ile	Thr	
		1360)				1365	5				1370)			
aag (ccg	ccc	tca	gac	gac	agc	ccg	gcc	cac	agc	agt	gcc	atc	ggg	ccc	4237
Lys I	Pro	Pro	Ser	Asp	Asp	Ser	Pro	Ala	His	Ser	Ser	Ala	Ile	Gly	Pro	
-	1375	5				1380)				1385	5				
gtc a	att	ggc	atc	atc	ctc	tct	ctc	ttc	gtc	atg	ggt	ggt	gtc	tat	ttt	4285
Val :	Ile	Gly	Ile	Ile	Leu	Ser	Leu	Phe	Val	Met	Gly	Gly	Val	Tyr	Phe	
1390					1395	5				1400)				1405	
gtg t	tgc	cag	cgc	gtg	gtg	tgc	cag	cgc	tat	gcg	999	gcc	aac	9 99	ccc	4333
Val (Cys	Gln	Arg	Val	Val	Cys	Gln	Arg	Tyr	Ala	Gly	Ala	Asn	Gly	Pro	
				1410)				1415	5				1420)	
ttc d	ccg	cac	gag	tat	gtc	agc	9 99	acc	ccg	cac	gtg	ccc	ctc	aat	ttc	4381
Phe I	Pro	His	Glu	Tyr	Val	Ser	Gly	Thr	Pro	His	Val	Pro	Leu	Asn	Phe	
			1425	5				1430)				1435	5		

1585	1590	1595
agg agc tac ttc cat ctc ttc	ccg ccc cct ccg tcc cc	c tgc acg gac 4909
Arg Ser Tyr Phe His Leu Phe	Pro Pro Pro Pro Ser Pr	o Cys Thr Asp
1600	1605 16	10
tca tcc tgacctcggc cgggccact	c tggcttctct gtgcccctg	t aaatagtttt 4965
Ser Ser		
1615		
aaatatgaac aaagaaaaaa atatatt	tta tgatttaaaa aataaat	ata attgggattt 5025
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gccggacaac atg gag gca gcg co	g ccc ggg ccg ccg tgg	ccg ctg ctg 109
Met Glu Ala Ala Pr	o Pro Gly Pro Pro Trp	Pro Leu Leu
1 5	10	
ctg ctg ctg ctg ctg ctg g	eg ctg tgc ggc tgc ccg	gcc ccc gcc 157
Leu Leu Leu Leu Leu Leu A	la Leu Cys Gly Cys Pro	Ala Pro Ala
15 20	25	
gcg gcc tcg ccg ctc ctg cta t	tt gcc aac cgc cgg gac	gta cgg ctg 205
Ala Ala Ser Pro Leu Leu Leu P	he Ala Asn Arg Arg Asp	Val Arg Leu
30 35	40	45
gtg gac gcc ggc gga gtc aag c	tg gag tcc acc atc gtg	gtc agc ggc 253
Val Asp Ala Gly Gly Val Lys L	eu Glu Ser Thr Ile Val	Val Ser Gly

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ctg gag gat gcg gcc gca gtg gac ttc cag ttt tcc aag gga gcc gtg Leu Glu Asp Ala Ala Ala Val Asp Phe Gln Phe Ser Lys Gly Ala Val tac tgg aca gac gtg agc gag gcc atc aag cag acc tac ctg aac Tyr Trp Thr Asp Val Ser Glu Glu Ala Ile Lys Gln Thr Tyr Leu Asn cag acg ggg gcc gcc gtg cag aac gtg gtc atc tcc ggc ctg gtc tct Gln Thr Gly Ala Ala Val Gln Asn Val Val Ile Ser Gly Leu Val Ser ccc gac ggc ctc gcc tgc gac tgg gtg ggc aag aag ctg tac tgg acg Pro Asp Gly Leu Ala Cys Asp Trp Val Gly Lys Lys Leu Tyr Trp Thr gac toa gag acc aac cgc atc gag gtg gcc aac ctc aat ggc aca tcc Asp Ser Glu Thr Asn Arg Ile Glu Val Ala Asn Leu Asn Gly Thr Ser egg aag gtg ete tte tgg eag gae ett gae eag eeg agg gee ate gee Arg Lys Val Leu Phe Trp Gln Asp Leu Asp Gln Pro Arg Ala Ile Ala ttg gac ccc gct cac ggg tac atg tac tgg aca gac tgg gtt gag acg Leu Asp Pro Ala His Gly Tyr Met Tyr Trp Thr Asp Trp Val Glu Thr ccc cgg att gag cgg gca ggg atg gat ggc agc acc cgg aag atc att Pro Arg Ile Glu Arg Ala Gly Met Asp Gly Ser Thr Arg Lys Ile Ile gtg gac tcg gac att tac tgg ccc aat gga ctg acc atc gac ctg gag Val Asp Ser Asp Ile Tyr Trp Pro Asn Gly Leu Thr Ile Asp Leu Glu gag cag aag ctc tac tgg gct gac gcc aag ctc agc ttc atc cac cgt

The first property of the second of the seco

Gli	ı Glr	n Tave	i T.e.i	j Tv2>	- Trr	, <u>∆</u>] ¬	Acr	. או		, T	. 0-:	o 151-	, T7		s Arg		
01.	. 011	y.	, nec			AIG	. АБР	ALO			ı Sei	: Pne	€ 11€				
				210					215					220			
															c ctg	781	L
Alá	Asr	ı Lei	ı Asp	o Gly	Ser	Phe	Arg	Gln	Lys	Va]	. Val	Glu	ı Gly	Sei	r Leu		
			225	5				230					235				
acc	cac	ccc	ttc	gcc	ctg	acg	ctc	tcc	a aa	gac	act	ctg	, tac	tgg	g aca	829	;
Thr	His	Pro	Phe	: Ala	Leu	Thr	Leu	Ser	Gly	Asp	Thr	Leu	Tyr	Trp	Thr		
		240	ı				245					250					
gac	tgg	cag	acc	cgc	tcc	atc	cat	gcc	tgc	aac	aag	cgc	act	gg 9	999	877	
Asp	Trp	Gln	Thr	Arg	Ser	Ile	His	Ala	Cys	Asn	Lys	Arg	Thr	Gly	Gly		
	255					260					265						
aag	agg	aag	gag	atc	ctg	agt	gcc	ctc	tac	tca	ccc	atg	gac	atc	cag	925	
Lys	Arg	Lys	Glu	Ile	Leu	Ser	Ala	Leu	Tyr	Ser	Pro	Met	Asp	Ile	Gln		
270					275					280					285		
gtg	ctg	agc	cag	gag	cgg	cag	cct	ttc	ttc	cac	act	cqc	tgt	gag	gag	973	
													Cys			5 / 5	
				290					295			5	-1 -	300			
gac	aat	ggc	qqc	tac	tcc	cac	cta	tac		cta	tcc	cca	agc		cct	1.00	-
													Ser			102	_
•		1	305	-1-					пси	цсu	261	PIO		GIU	PIO		
ttc	tac	202		77	t ~ ~			310					315				
													gac			1069)
riie	ıyı		Cys	Ala	Cys			GIY	Val	Gln	Leu	Gln	Asp	Asn	Gly		
		320					325					330					
													gcc			1117	,
Arg	Thr	Cys	Lys	Ala	Gly	Ala	Glu	Glu	Val	Leu	Leu	Leu	Ala	Arg	Arg		
	335					340					345						
acg	gac	cta	cgg	agg	atc	tcg	ctg (gac	acg	ccg	gac	ttc	acc	gac	atc	1165	i
Chr	Asp	Leu	Arg	Arg	Ile	Ser :	Leu 1	Asp	Thr	Pro	Asp	Phe	Thr	Asp	Ile		
350					355					360					365		

gtg	ctg	cag	gtg	gac	gac	atc	cgg	cac	gcc	att	gcc	atc	gac	tac	gac	1213
Val	Leu	Gln	Val	Asp	Asp	Ile	Arg	His	Ala	Ile	Ala	Ile	Asp	Tyr	Asp	
				370					375					380		
ccg	cta	gag	ggc	tat	gtc	tac	tgg	aca	gat	gac	gag	gtg	cgg	gcc	atc	1261
Pro	Leu	Glu	Gly	Tyr	Val	Tyr	Trp	Thr	Asp	Asp	Glu	Val	Arg	Ala	Ile	
			385					390					395			
cgc	agg	gcg	tac	ctg	gac	ggg	tct	aaa	gcg	cag	acg	ctg	gtc	aac	acc	1309
Arg	Arg	Ala	Tyr	Leu	Asp	Gly	Ser	Gly	Ala	Gln	Thr	Leu	Val	Asn	Thr	
		400					405					410				
gag	atc	aac	gac	ccc	gat	ggc	atc	gcg	gtc	gac	tgg	gtg	gcc	cga	aac	1357
Glu	Ile	Asn	Asp	Pro	Asp	Gly	Ile	Ala	Val	Asp	Trp	Val	Ala	Arg	Asn	
	415					420					425					
ctc	tac	tgg	acc	gac	acg	ggc	acg	gac	cgc	atc	gag	gtg	acg	cgc	ctc	1405
Leu	Tyr	Trp	Thr	Asp	Thr	Gly	Thr	Asp	Arg	Ile	Glu	Val	Thr	Arg	Leu	
430					435					440					445	
aac	ggc	acc	tcc	cgc	aag	atc	ctg	gtg	tcg	gag	gac	ctg	gac	gag	CCC	1453
Asn	Gly	Thr	Ser	Arg	Lys	Ile	Leu	Val	Ser	Glu	Asp	Leu	Asp	Glu	Pro	
				450					455					460		
cga	gcc	atc	gca	ctg	cac	CCC	gtg	atg	ggc	ctc	atg	tac	tgg	aca	gac	1501
Arg	Ala	Ile	Ala	Leu	His	Pro	Val	Met	Gly	Leu	Met	Tyr	Trp	Thr	Asp	
			465					470					475			
tgg	gga	gag	aac	cct	aaa	atc	gag	tgt	gcc	aac	ttg	gat	9 99	cag	gag	1549
Trp	Gly	Glu	Asn	Pro	Lys	Ile	Glu	Cys	Ala	Asn	Leu	Asp	Gly	Gln	Glu	
		480					485					490				
	_	gtg	-	_												1597
Arg	_	Val	Leu	Val	Asn	Ala	Ser	Leu	Gly	Trp	Pro	Asn	Gly	Leu	Ala	
	495					500					505					
		ctg														1645
Leu	Asp	Leu	Gln	Glu	Gly	Lys	Leu	Tyr	Trp	Gly	Asp	Ala	Lys	Thr	Asp	

510					515					520					525	
aag	atc	gag	gtg	atc	aat	gtt	gat	aaa	acg	aag	agg	cgg	acc	ctc	ctg	1693
Lys	Ile	Glu	Val	Ile	Asn	Val	Asp	Gly	Thr	Lys	Arg	Arg	Thr	Leu	Leu	
				530					535					540		
gag	gac	aag	ctc	ccg	cac	att	ttc	ggg	ttc	acg	ctg	ctg	999	gac	ttc	1741
Glu	Asp	Lys	Leu	Pro	His	Ile	Phe	Gly	Phe	Thr	Leu	Leu	Gly	Asp	Phe	
			545					550					555			
atc	tac	tgg	act	gac	tgg	cag	cgc	cgc	agc	atc	gag	cgg	gtg	cac	aag	1789
Ile	Tyr	Trp	Thr	Asp	Trp	Gln	Arg	Arg	Ser	Ile	Glu	Arg	Val	His	Lys	
		560					565					570				
gtc	aag	gcc	agc	cgg	gac	gtc	atc	att	gac	cag	ctg	ccc	gac	ctg	atg	1837
Val	Lys	Ala	Ser	Arg	Asp	Val	Ile	Ile	Asp	Gln	Leu	Pro	Asp	Leu	Met	
	575					580					585					
999	ctc	aaa	gct	gtg	aat	gtg	gcc	aag	gtc	gtc	gga	acc	aac	ccg	tgt	1885
Gly	Leu	Lys	Ala	Val	Asn	Val	Ala	Lys	Val	Val	Gly	Thr	Asn	Pro	Cys	
590					595					600					605	
gcg	gac	agg	aac	ggg	aaa	tgc	agc	cac	ctg	tgc	ttc	ttc	aca	ccc	cac	1933
Ala	Asp	Arg	Asn	Gly	Gly	Cys	Ser	His	Leu	Cys	Phe	Phe	Thr	Pro	His	
				610					615					620		
gca	acc	cgg	tgt	ggc	tgc	ccc	atc	ggc	ctg	gag	ctg	ctg	agt	gac	atg	1981
Ala	Thr	Arg	Cys	Gly	Cys	Pro	Ile	Gly	Leu	Glu	Leu	Leu	Ser	Asp	Met	
			625					630					635			
aag	acc	tgc	atc	gtg	cct	gag	gcc	ttc	ttg	gtc	ttc	acc	agc	aga	gcc	2029
Lys	Thr	Cys	Ile	Val	Pro	Glu	Ala	Phe	Leu	Val	Phe	Thr	Ser	Arg	Ala	
		640					645					650				
gcc	atc	cac	agg	atc	tcc	ctc	gag	acc	aat	aac	aac	gac	gtg	gcc	atc	2077
Ala	Ile	His	Arg	Ile	Ser	Leu	Glu	Thr	Asn	Asn	Asn	Asp	Val	Ala	Ile	
	655					660					665					
ccg	ctc	acg	ggc	gtc	aag	gag	gcc	tca	gcc	ctg	gac	ttt	gat	gtg	tcc	2125

Pro	Leu	Thr	Gly	Val	Lys	Glu	Ala	Ser	Ala	Leu	Asp	Phe	Asp	Val	Ser	
670					675					680					685	
aac	aac	cac	atc	tac	tgg	aca	gac	gtc	agc	ctg	aag	acc	atc	agc	cgc	2173
Asn	Asn	His	Ile	Tyr	Trp	Thr	Asp	Val	Ser	Leu	Lys	Thr	Ile	Ser	Arg	
				690					695					700		
gcc	ttc	atg	aac	ggg	agc	tcg	gtg	gag	cac	gtg	gtg	gag	ttt	ggc	ctt	2221
Ala	Phe	Met	Asn	Gly	Ser	Ser	Val	Glu	His	Val	Val	Glu	Phe	Gly	Leu	
			705					710					715			
gac	tac	ccc	gag	ggc	atg	gcc	gtt	gac	tgg	atg	ggc	aag	aac	ctc	tac	2269
Asp	Tyr	Pro	Glu	Gly	Met	Ala	Val	Asp	Trp	Met	Gly	Lys	Asn	Leu	Tyr	
		720					725					730				
tgg	gcc	gac	act	999	acc	aac	aga	atc	gaa	gtg	gcg	cgg	ctg	gac	a aa	2317
Trp	Ala	Asp	Thr	Gly	Thr	Asn	Arg	Ile	Glu	Val	Ala	Arg	Leu	Asp	Gly	
	735					740					745					
cag	ttc	cgg	caa	gtc	ctc	gtg	tgg	agg	gac	ttg	gac	aac	ccg	agg	tcg	2365
Gln	Phe	Arg	Gln	Val	Leu	Val	Trp	Arg	Asp	Leu	Asp	Asn	Pro	Arg	Ser	
750					755					760					765	
ctg	gcc	ctg	gat	ccc	acc	aag	ggc	tac	atc	tac	tgg	acc	gag	tgg	ggc	2413
Leu	Ala	Leu	Asp	Pro	Thr	Lys	Gly	Tyr	Ile	Tyr	Trp	Thr	Glu	Trp	Gly	
				770					775					780		
ggc	aag	ccg	agg	atc	gtg	cgg	gcc	ttc	atg	gac	999	acc	aac	tgc	atg	2461
Gly	Lys	Pro	Arg	Ile	Val	Arg	Ala	Phe	Met	Asp	Gly	Thr	Asn	Cys	Met	
			785					790					795			
acg	ctg	gtg	gac	aag	gtg	ggc	cgg	gcc	aac	gac	ctc	acc	att	gac	tac	2509
Thr	Leu	Val	Asp	Lys	Val	Gly	Arg	Ala	Asn	Asp	Leu	Thr	Ile	Asp	Tyr	
		800					805					810				
gct	gac	cag	cgc	ctc	tac	tgg	acc	gac	ctg	gac	acc	aac	atg	atc	gag	2557
Ala	Asp	Gln	Arg	Leu	Tyr	Trp	Thr	Asp	Leu	Asp	Thr	Asn	Met	Ile	Glu	
	815					820					825					

tcg	tcc	aac	atg	ctg	ggt	cag	gag	cgg	gtc	gtg	att	gcc	gac	gat	ctc	2605
Ser	Ser	Asn	Met	Leu	Gly	Gln	Glu	Arg	Val	Val	Ile	Ala	Asp	Asp	Leu	
830					835					840					845	
ccg	cac	ccg	ttc	ggt	ctg	acg	cag	tac	agc	gat	tat	atc	tac	tgg	aca	2653
Pro	His	Pro	Phe	Gly	Leu	Thr	Gln	Tyr	Ser	Asp	Tyr	Ile	Tyr	Trp	Thr	
				850					855					860		
gac	tgg	aat	ctg	cac	agc	att	gag	cgg	gcc	gac	aag	act	agc	ggc	cgg	2701
Asp	Trp	Asn	Leu	His	Ser	Ile	Glu	Arg	Ala	Asp	Lys	Thr	Ser	Gly	Arg	
			865					870					875			
aac	cgc	acc	ctc	atc	cag	ggc	cac	ctg	gac	ttc	gtg	atg	gac	atc	ctg	2749
Asn	Arg	Thr	Leu	Ile	Gln	Gly	His	Leu	Asp	Phe	Val	Met	Asp	Ile	Leu	
		880					885					890				
gtg	ttc	cac	tcc	tcc	cgc	cag	gat	ggc	ctc	aat	gac	tgt	atg	cac	aac	2797
Val	Phe	His	Ser	Ser	Arg	Gln	Asp	Gly	Leu	Asn	Asp	Cys	Met	His	Asn	
	895					900					905					
aac	999	cag	tgt	999	cag	ctg	tgc	ctt	gcc	atc	ccc	ggc	ggc	cac	cgc	2845
Asn	Gly	Gln	Cys	Gly	Gln	Leu	Cys	Leu	Ala	Ile	Pro	Gly	Gly	His	Arg	
910					915					920					925	
tgc	ggc	tgc	gcc	tca	cac	tac	acc	ctg	gac	CCC	agc	agc	cgc	aac	tgc	2893
Cys	Gly	Cys	Ala	Ser	His	Tyr	Thr	Leu	Asp	Pro	Ser	Ser	Arg	Asn	Cys	
				930					935					940		
agc	ccg	CCC	acc	acc	ttc	ttg	ctg	ttc	agc	cag	aaa	tct	gcc	atc	agt	2941
Ser	Pro	Pro	Thr	Thr	Phe	Leu	Leu	Phe	Ser	Gln	Lys	Ser	Ala	Ile	Ser	
			945					950					955			
cgg	atg	atc	ccg	gac	gac	cag	cac	agc	ccg	gat	ctc	atc	ctg	ccc	ctg	2989
Arg	Met	Ile	Pro	Asp	Asp	Gln	His	Ser	Pro	Asp	Leu	Ile	Leu	Pro	Leu	
		960					965					970				
cat	gga	ctg	agg	aac	gtc	aaa	gcc	atc	gac	tat	gac	cca	ctg	gac	aag	3037
His	Gly	Leu	Arg	Asn	Val	Lys	Ala	Ile	Asp	Tyr	Asp	Pro	Leu	Asp	Lys	

975	980	985		
ttc atc tac tgg gtg	g gat ggg cgc c	ag aac atc aag cga	ı gcc aag gac 308	35
Phe Ile Tyr Trp Val	l Asp Gly Arg G	ln Asn Ile Lys Arc	Ala Lys Asp	
990	995	1000	1005	
gac ggg acc cag cco	c ttt gtt ttg a	cc tct ctg agc caa	ggc caa aac 313	3
Asp Gly Thr Gln Pro	o Phe Val Leu Th	ır Ser Leu Ser Glr	. Gly Gln Asn	
101	10	1015	1020	
cca gac agg cag ccc	cac gac ctc ag	gc atc gac atc tac	agc cgg aca 318	1
Pro Asp Arg Gln Pro	His Asp Leu Se	er Ile Asp Ile Tyr	Ser Arg Thr	
1025	10	30	1035	
ctg ttc tgg acg tgc	gag gcc acc aa	it acc atc aac gtc	cac agg ctg 322	9
Leu Phe Trp Thr Cys	Glu Ala Thr As	n Thr Ile Asn Val	His Arg Leu	
1040	1045	105	0	
agc ggg gaa gcc atg	g ggg gtg gtg ct	g cgt ggg gac cgc	gac aag ccc 327	7
Ser Gly Glu Ala Met	: Gly Val Val Le	u Arg Gly Asp Arg	Asp Lys Pro	
1055	1060	1065		
agg gcc atc gtc gtc	aac gcg gag cg	a ggg tac ctg tac	ttc acc aac 332	5
Arg Ala Ile Val Val	Asn Ala Glu Ar	g Gly Tyr Leu Tyr	Phe Thr Asn	
1070	1075	1080	1085	
atg cag gac cgg gca	gcc aag atc ga	a cgc gca gcc ctg	gac ggc acc 3373	3
Met Gln Asp Arg Ala	Ala Lys Ile Gl	u Arg Ala Ala Leu	Asp Gly Thr	
109	0	1095	1100	
gag cgc gag gtc ctc	ttc acc acc gg	c ctc atc cgc cct	gtg gcc ctg 3423	L
Glu Arg Glu Val Leu	Phe Thr Thr Gl	y Leu Ile Arg Pro	Val Ala Leu	
1105	11	10	1115	
gtg gtg gac aac aca	ctg ggc aag ctg	g ttc tgg gtg gac	gcg gac ctg 3469)
Val Val Asp Asn Thr	Leu Gly Lys Le	1 Phe Trp Val Asp	Ala Asp Leu	
1120	1125	1130		
aag cgc att gag agc	tgt gac ctg tca	a ggg gcc aac cgc	ctg acc ctg 3517	,

Lys Arg Ile Glu Ser Cys Asp Leu Ser Gly Ala Asn Arg Leu Thr Leu gag gac gcc aac atc gtg cag cct ctg ggc ctg acc atc ctt ggc aag Glu Asp Ala Asn Ile Val Gln Pro Leu Gly Leu Thr Ile Leu Gly Lys cat ctc tac tgg atc gac cgc cag cag cag atg atc gag cgt gtg gag His Leu Tyr Trp Ile Asp Arg Gln Gln Gln Met Ile Glu Arg Val Glu aag acc acc ggg gac aag cgg act cgc atc cag ggc cgt gtc gcc cac Lys Thr Thr Gly Asp Lys Arg Thr Arg Ile Gln Gly Arg Val Ala His ctc act ggc atc cat gca gtg gag gaa gtc agc ctg gag gag ttc tca Leu Thr Gly Ile His Ala Val Glu Glu Val Ser Leu Glu Glu Phe Ser gee cae cea tgt gee cgt gae aat ggt gge tge tee cae ate tgt att Ala His Pro Cys Ala Arg Asp Asn Gly Gly Cys Ser His Ile Cys Ile gcc aag ggt gat ggg aca cca cgg tgc tca tgc cca gtc cac ctc gtg Ala Lys Gly Asp Gly Thr Pro Arg Cys Ser Cys Pro Val His Leu Val ctc ctg cag aac ctg ctg acc tgt gga gag ccg ccc acc tgc tcc ccg Leu Leu Gln Asn Leu Leu Thr Cys Gly Glu Pro Pro Thr Cys Ser Pro gac cag ttt gca tgt gcc aca ggg gag atc gac tgt atc ccc ggg gcc Asp Gln Phe Ala Cys Ala Thr Gly Glu Ile Asp Cys Ile Pro Gly Ala tgg cgc tgt gac ggc ttt ccc gag tgc gat gac cag agc gac gag Trp Arg Cys Asp Gly Phe Pro Glu Cys Asp Asp Gln Ser Asp Glu Glu

ggc	tgo	CCC	gtg	g tgo	t tcc	gcc	gcc	cag	, ttc	ccc	: tg	g gcg	g cgg	g ggt	cag	3997	
Gly	Cys	Pro	o Val	Cys	s Ser	Ala	Ala	Gln	Phe	Pro	Суз	s Ala	a Arg	g Gly	Gln Gln		
	129	5				130	0				130	5					
tgt	gtg	gad	ctg	g	ctg	cgc	tgc	gac	ggc	gag	gca	gad	tgt:	cag	gac	4045	
Cys	Val	Asp) Leu	Arg	, Leu	Arg	Cys	Asp	Gly	Glu	Ala	. Asp	Суз	Gln	Asp		
131	0				131	5				132	0				1325		
cgc	tca	gac	gag	gtg	gac	tgt	gac	gcc	atc	tgc	ctg	ccc	aac	cag	ttc	4093	
Arg	Ser	Asp	Glu	Val	Asp	Cys	Asp	Ala	Ile	Cys	Leu	Pro	Asn	. Gln	Phe		
				133	0				133!	5				134	0		
cgg	tgt	gcg	agc	ggc	cag	tgt	gtc	ctc	atc	aaa	cag	cag	tgc	gac	tcc	4141	
					Gln												
			134					135					135				
ttc	ccc	gac	tgt	atc	gac	ggc	tcc	gac	gag	ctc	atg	tqt	qaa	atc	acc	4189	
					Asp												
		136					1365					137					
aag	ccg	ccc	tca	gac	gac	agc	ccq	gcc	cac	agc	agt	acc	atc	aaa	CCC	4237	
					Asp											100,	
	1375			_	-	1380					138			017			
gtc	att	aac	atc	atc	ctc			ttc	atc	ato			ata	t a t		4205	
					Leu											4285	
1390		1			1395		Dea	1110	vai	1400		GIY	vai	ıyı			
		cac	cac	ata											1405		
gtg Val																4333	
Val	СуБ	0111	Arg			Cys	GIII			Ala	GTÀ	Ala	Asn				
				1410					1415					1420			
ttc	ccg	cac	gag	tat	gtc	agc	a aa	acc	ccg	cac	gtg	ccc	ctc	aat	ttc	4381	
Phe 1	Pro	His	Glu	Tyr	Val	Ser	Gly	Thr	Pro :	His	Val	Pro	Leu	Asn	Phe		
			1425					1430					1435				
ata g	gcc	ccg	ggc	ggt	tcc	cag	cat (ggc	ccc	ttc	aca	ggc	atc	gca	tgc	4429	
Ile A	Ala .	Pro	Gly	Gly	Ser	Gln :	His (Gly	Pro 1	Phe	Thr	Gly	Ile	Ala	Cys		

			•				244	_				3.45	•			
		1440)				1445)				145	U			
gga	aag	tcc	atg	atg	agc	tcc	gtg	agc	ctg	atg	9 99	ggc	cgg	ggc	999	4477
Gly	Lys	Ser	Met	Met	Ser	Ser	Val	Ser	Leu	Met	Gly	Gly	Arg	Gly	Gly	
	1455	5				1460)				1465	5				
gtg	ccc	ctc	tac	gac	cgg	aac	cac	gtc	aca	999	gcc	tcg	tcc	agc	agc	4525
Val	Pro	Leu	Tyr	Asp	Arg	Asn	His	Val	Thr	Gly	Ala	Ser	Ser	Ser	Ser	
1470)				1475	5				1480)				1485	
tcg	tcc	agc	acg	aag	gcc	acg	ctg	tac	ccg	ccg	atc	ctg	aac	ccg	ccg	4573
Ser	Ser	Ser	Thr	Lys	Ala	Thr	Leu	Tyr	Pro	Pro	Ile	Leu	Asn	Pro	Pro	
				1490)				1499	5				150	0	
CCC	tcc	cca	acc	aca	gac	CCC	tcc	cta	tac	aac	atq	gac	atq	ttc	tac	4621
		_	_			Pro										1001
FIO	561	PIO	1505		App	FIQ	561	1510		ASII	Mec	Asp	1515		1 7 1	
	.									.						4660
						act										4669
Ser	Ser	Asn	Ile	Pro	Ala	Thr	Ala	Arg	Pro	Tyr	Arg	Pro	Tyr	Ile	Ile	
		1520)				1525	5				1530)			
cga	gga	atg	gcg	ccc	ccg	acg	acg	ccc	tgc	agc	acc	gac	gtg	tgt	gac	4717
Arg	Gly	Met	Ala	Pro	Pro	Thr	Thr	Pro	Cys	Ser	Thr	Asp	Val	Cys	Asp	
	1535	5				1540)				1545	5				
agc	gac	tac	agc	gcc	agc	cgc	tgg	aag	gcc	agc	aag	tac	tac	ctg	gat	4765
Ser	Asp	Tyr	Ser	Ala	Ser	Arg	Trp	Lys	Ala	Ser	Lys	Tyr	Tyr	Leu	Asp	
1550)				1555	5				1560)				1565	
ttg	aac	tcg	gac	tca	gac	ccc	tat	сса	ccc	cca	ccc	acg	ccc	cac	agc	4813
Leu	Asn	Ser	Asp	Ser	Asp	Pro	Tyr	Pro	Pro	Pro	Pro	Thr	Pro	His	Ser	
				1570)				1575	5				1580)	
cag	tac	ctg	tcg	gcg	gag	gac	agc	tgc	ccg	CCC	tcg	ccc	gcc	acc	gag	4861
Gln	Tyr	Leu	Ser	Ala	Glu	Asp	Ser	Cys	Pro	Pro	Ser	Pro	Ala	Thr	Glu	
	-		1585			_		1590					1595			
acc	acc	tac			ctc	ttc	cca			ככמ	tcc	ccc			gac	4909
~99	age	cac		Cac			229			229			-gc	ucy	J~C	

Arg Ser Tyr Phe His Leu Phe Pro Pro Pro Pro Ser Pro Cys Thr Asp	
1600 1605 1610	
tca tcc tgacctcggc cgggccactc tggcttctct gtgcccctgt aaatagtttt	4965
Ser Ser	
1615	
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Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30	
Leu Leu Leu Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala	
Leu Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala 35 40 45	
Leu Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala 35 40 45 Gly Gly Val Lys Leu Glu Ser Thr Ile Val Val Ser Gly Leu Glu Asp	
Leu Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala 35 40 45 Gly Gly Val Lys Leu Glu Ser Thr Ile Val Val Ser Gly Leu Glu Asp 50 55 60	
Leu Leu Leu Leu Ala Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala 35 40 45 Gly Gly Val Lys Leu Glu Ser Thr Ile Val Val Ser Gly Leu Glu Asp 50 55 60 Ala Ala Ala Val Asp Phe Gln Phe Ser Lys Gly Ala Val Tyr Trp Thr	
Leu Leu Leu Leu Leu Cys Gly Cys Pro Ala Pro Ala Ala Ala Ala Ser 20 25 30 Pro Leu Leu Leu Leu Phe Ala Asn Arg Arg Asp Val Arg Leu Val Asp Ala 35 40 45 Gly Gly Val Lys Leu Glu Ser Thr Ile Val Val Ser Gly Leu Glu Asp 50 55 60 Ala Ala Ala Val Asp Phe Gln Phe Ser Lys Gly Ala Val Tyr Trp Thr 65 70 75 80	

Leu	ı Ala	. Cys	Asp	Trp	Val	Gly	Lys	Lys	Leu	туг	Trp	Thr	Asp	Ser	Glu
		115					120					125	i		
Thr	Asn	Arg	Ile	Glu	Val	Ala	Asn	Leu	Asn	Gly	Thr	Ser	Arg	Lys	Va]
	130					135					140				
Leu	Phe	Trp	Gln	Asp	Leu	Asp	Gln	Pro	Lys	Ala	Ile	Ala	. Leu	Asp	Pro
145					150					155	i				160
Ala	His	Gly	Tyr	Met	Tyr	Trp	Thr	Asp	Trp	Gly	Glu	Thr	Pro	Arg	Ile
				165					170					175	
Glu	Arg	Ala	Gly	Met	Asp	Gly	Ser	Thr	Arg	Lys	Ile	Ile	Val	Asp	Ser
			180					185					190		
Asp	Ile	Tyr	Trp	Pro	Asn	Gly	Leu	Thr	Ile	Asp	Leu	Glu	Glu	Gln	Lys
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Leu	Tyr	Trp	Ala	Asp	Ala	Lys	Leu	Ser	Phe	Ile	His	Arg	Ala	Asn	Leu
	210					215					220				
Asp	Gly	Ser	Phe	Arg	Gln	Lys	Val	Val	Glu	Gly	Ser	Leu	Thr	His	Pro
225					230					235					240
Phe	Ala	Leu	Thr	Leu	Ser	Gly	Asp	Thr	Leu	Tyr	Trp	Thr	Asp	Trp	Gln
				245					250					255	
Thr	Arg	Ser	Ile	His	Ala	Cys	Asn	Lys	Arg	Thr	Gly	Gly	Lys	Arg	Lys
			260					265					270		
Glu	Ile	Leu	Ser	Ala	Leu	Tyr	Ser	Pro	Met	Asp	Ile	Gln	Val	Leu	Ser
		275					280					285			
Gln	Glu	Arg	Gln	Pro	Phe	Phe	His	Thr	Arg	Cys	Glu	Glu	Asp	Asn	Gly
	290					295					300				
Gly	Trp	Ser	His	Leu	Cys	Leu	Leu	Ser	Pro	Ser	Glu	Pro	Phe	Tyr	Thr
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Cys	Ala	Cys	Pro	Thr	Gly	Val	Gln	Met	Gln	Asp	Asn	Gly	Arg	Thr	Cys
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Lys	Ala	Gly	Ala	Glu	Glu	Val	Leu	Leu	Leu	Ala	Arg	Arg	Thr	Asp	Leu

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Tyr	Leu	Asp	Gly	Ser	Gly	Ala	Gln	Thr	Leu	Val	Asn	Thr	Glu	Ile	Asn
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Asp	Pro	Asp	Gly	Ile	Ala	Val	Asp	Trp	Val	Ala	Arg	Asn	Leu	Tyr	Trp
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Thr	Asp	Thr	Gly	Thr	Asp	Arg	Ile	Glu	Val	Thr	Arg	Leu	Asn	Gly	Thr
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Ser	Arg	Lys	Ile	Leu	Val	Ser	Glu	Asp	Leu	Asp	Glu	Pro	Arg	Ala	Ile
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Ala	Leu	His	Pro	Val	Met	Gly	Leu	Met	Tyr	Trp	Thr	Asp	Trp	Gly	Glu
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Asn	Pro	Lys	Ile	Glu	Cys	Ala	Asn	Leu	Asp	Gly	Gln	Glu	Arg	Arg	Val
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Leu	Val	Asn	Ala	Ser	Leu	Gly	Trp	Pro	Asn	Gly	Leu	Ala	Leu	Asp	Leu
			500					505					510		
Gln	Glu	Gly	Lys	Leu	Tyr	Trp	Gly	Asp	Ala	Lys	Thr	Asp	Lys	Ile	Glu
		515					520					525			
Val	Ile	Asn	Val	Asp	Gly	Thr	Lys	Arg	Arg	Thr	Leu	Leu	Glu	Asp	Lys
	530					535					540				
Leu	Pro	His	Ile	Phe	Gly	Phe	Thr	Leu	Leu	Gly	Asp	Phe	Ile	Tyr	Trp
545					550					555					560
Thr	Asp	Trp	Gln	Arg	Arg	Ser	Ile	Glu	Arg	Val	His	Lys	Val	Lys	Ala
				565					570					575	

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Ser	Arg	Asp	Val	Ile	Ile	Asp	Gln	Leu	Pro	Asp	Leu	Met	Gly	Leu	Lys
			580					585					590		
Ala	. Val	Asn	Val	Ala	Lys	Val	Val	Gly	Thr	Asn	Pro	Cys	Ala	Asp	Arg
		595					600					605			
Asn	Gly	Gly	Cys	Ser	His	Leu	Cys	Phe	Phe	Thr	Pro	His	Ala	Thr	Arg
	610					615					620				
Cys	Gly	Cys	Pro	Ile	Gly	Leu	Glu	Leu	Leu	Ser	Asp	Met	Lys	Thr	Cys
625					630					635					640
Ile	Val	Pro	Glu	Ala	Phe	Leu	Val	Phe	Thr	Ser	Arg	Ala	Ala	Ile	His
				645					650					655	
Arg	Ile	Ser	Leu	Glu	Thr	Asn	Asn	Asn	Asp	Val	Ala	Ile	Pro	Leu	Thr
			660					665					670		
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		675					680					685			
Ile	Tyr	Trp	Thr	Asp	Val	Ser	Leu	Lys	Asn	Ile	Ser	Arg	Ala	Phe	Met
	690					695					700				
Asn	Gly	Ser	Ser	Val	Glu	His	Val	Val	Glu	Phe	Gly	Leu	Asp	Tyr	Pro
705					710					715					720
Glu	Gly	Met	Ala	Val	Asp	Trp	Met	Gly	Lys	Asn	Leu	Tyr	Trp	Ala	Asp
				725					730					735	
Thr	Gly	Thr	Asn	Arg	Ile	Glu	Val	Ala	Arg	Leu	Asp	Gly	Gln	Phe	Arg
			740					745					750		
Gln	Val	Leu	Val	Trp	Arg	Asp	Leu	Asp	Asn	Pro	Arg	Ser	Leu	Ala	Leu
		755					760					765			
Asp	Pro	Thr	Lys	Gly	Tyr	Ile	Tyr	Trp	Thr	Glu	Trp	Gly	Gly	Lys	Pro
	770					775					780				
Arg	Ile	Val	Arg	Ala	Phe	Met	Asp	Gly	Thr	Asn	Cys	Met	Thr	Leu	Val
785					790					795					800
Asp	Lvs	Val	Glv	Ara	Ala	Asn	Asp	Leu	Thr	Tle	Δen	ጥኒ/ዮ	בומ	λαn	Gln

Arg Leu Tyr Trp Thr Asp Leu Asp Thr Asn Met Ile Glu Ser Ser Asn

Met Leu Gly Gln Glu Arg Val Val Ile Ala Asp Asp Leu Pro His Pro

	Phe	Gly	Leu	Thr	Gln	Tyr	Ser	Asp	Tyr	· Ile	Tyr	Trp	Thr	Asp	Trp	Asn
		850					855					860				
	Leu	His	Ser	Ile	Glu	Arg	Ala	Asp	Lys	Thr	Ser	Gly	Arg	Asn	Arg	Thr
	865					870					875					880
	Leu	Ile	Gln	Gly	His	Leu	Asp	Phe	Val	Met	Asp	Ile	Leu	Val	Phe	His
					885					890					895	
100 100 100 100 100	Ser	Ser	Arg	Gln	Asp	Gly	Leu	Asn	Asp	Сув	Met	His	Asn	Asn	Gly	Gln
				900					905					910		
	Cys	Gly	Gln	Leu	Cys	Leu	Ala	Ile	Pro	Gly	Gly	His	Arg	Cys	Gly	Cys
			915					920					925			
i	Ala	Ser	His	Tyr	Thr	Leu	Asp	Pro	Ser	Ser	Arg	Asn	Cys	Ser	Pro	Pro
To the second se		930					935					940				
	Thr	Thr	Phe	Leu	Leu	Phe	Ser	Gln	Lys	Ser	Ala	Ile	Ser	Arg	Met	Ile
1	945					950					955					960
	Pro	Asp	Asp	Gln	His	Ser	Pro	Asp	Leu	Ile	Leu	Pro	Leu	His	Gly	Leu
					965					970					975	
	Arg	Asn	Val	Lys	Ala	Ile	Asp	Tyr	Asp	Pro	Leu	Asp	Lys	Phe	Ile	Tyr
				980					985					990		
	Trp	Val	Asp	Gly	Arg	Gln	Asn	Ile	Lys	Arg	Ala	Lys	Asp	Asp	Gly	Thr
			995					1000)				1005	;		
	Gln	Pro	Phe	Val	Leu	Thr	Ser	Leu	Ser	Gln	Gly	Gln	Asn	Pro	Asp	Arg

Gln Pro His Asp Leu Ser Ile Asp Ile Tyr Ser Arg Thr Leu Phe Trp

Thr	Cys	Glu	Ala	Thr	Asn	Thr	Ile	Asn	Val	His	Arg	Leu	Ser	Gly	Glu
				104	5				105	0				105	5
Ala	Met	Gly	Val	Val	Leu	Arg	Gly	Asp	Arg	Asp	Lys	Pro	Arg	Ala	Ile
			106	0				106	5				107	0	
Val	Val	Asn	Ala	Glu	Arg	Gly	Tyr	Leu	Tyr	Phe	Thr	Asn	Met	Gln	Asp
		107	5				108	0				108	5		
Arg	Ala	Ala	Lys	Ile	Glu	Arg	Ala	Ala	Leu	Asp	Gly	Thr	Glu	Arg	Glu
	109	0				109	5				110	0			
Val	Leu	Phe	Thr	Thr	Gly	Leu	Ile	Arg	Pro	Val	Ala	Leu	Val	Val	Asp
110	5				111	0				111	5				1120
Asn	Thr	Leu	Gly	Lys	Leu	Phe	Trp	Val	Asp	Ala	Asp	Leu	Lys	Arg	Ile
				112	5				113	0				113	5
Glu	Ser	Cys	Asp	Leu	Ser	Gly	Ala	Asn	Arg	Leu	Thr	Leu	Glu	Asp	Ala
			114	0				1145	5				115	0	
Asn	Ile	Val	Gln	Pro	Leu	Gly	Leu	Thr	Ile	Leu	Gly	Lys	His	Leu	Tyr
		115	5				1160)				1169	5		
Trp	Ile	Asp	Arg	Gln	Gln	Gln	Met	Ile	Glu	Arg	Val	Glu	Lys	Thr	Thr
	1170)				1175	5				1180)			
Gly	Asp	Lys	Arg	Thr	Arg	Ile	Gln	Gly	Arg	Val	Ala	His	Leu	Thr	Gly
118	5				1190)				1199	5				1200
Ile	His	Ala	Val	Glu	Glu	Val	Ser	Leu	Glu	Glu	Phe	Ser	Ala	His	Pro
				1205	5				1210)				1215	5
Cys	Ala	Arg	Asp	Asn	Gly	Gly	Cys	Ser	His	Ile	Cys	Ile	Ala	Lys	Gly
			1220)				1225					1230)	
Asp	Gly	Thr	Pro	Arg	Cys	Ser	Cys	Pro	Val	His	Leu	Val	Leu	Leu	Gln
		1235	i				1240)				1245			
Asn	Leu	Leu	Thr	Cys	Gly	Glu	Pro	Pro	Thr	Cys	Ser	Pro	Asp	Gln	Phe
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Ala	Cys	Ala	Thr	Gly	Glu	Ile	Asp	Cys	Ile	Pro	Gly	Ala	Trp	Arg	Cys

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Asp	Gly	Phe	Pro	Glu	Cys	Asp	Asp	Gln	Ser	Asp	Glu	Glu	Gly	Cys	Pro
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Val	Cys	Ser	Ala	Ala	Gln	Phe	Pro	Cys	Ala	Arg	Gly	Gln	Cys	Val	Asp
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Leu	Arg	Leu	Arg	Cys	Asp	Gly	Glu	Ala	Asp	Cys	Gln	Asp	Arg	Ser	Asp
		1315	ō				1320)				1329	5		
Glu	Val	Asp	Cys	Asp	Ala	Ile	Cys	Leu	Pro	Asn	Gln	Phe	Arg	Cys	Ala
	1330)				1335	5				1340)			
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Cys	Ile	Asp	Gly	Ser	Asp	Glu	Leu	Met	Cys	Glu	Ile	Thr	Lys	Pro	Pro
				1365	5				1370)				1375	5
Ser	Asp	Asp	Ser	Pro	Ala	His	Ser	Ser	Ala	Ile	Gly	Pro	Val	Ile	Gly
			1380)				1385	5				1390)	
Ile	Ile	Leu	Ser	Leu	Phe	Val	Met	Gly	Gly	Val	Tyr	Phe	Val	Cys	Gln
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		1395	5				1400)				1405	5	-	
Arg	Val			Gln	Arg	Tyr			Ala	Asn	Gly			Pro	His
Arg	Val	Val		Gln	Arg	Tyr 1415	Ala		Ala	Asn	Gly 1420	Pro		-	His
	1410	Val	Cys			1415	Ala	Gly			1420	Pro	Phe	-	
	1410 Tyr	Val	Cys			1415 Pro	Ala	Gly			1420 Asn	Pro	Phe	Pro	
Glu 1425	1410 Tyr	Val) Val	Cys Ser	Gly	Thr	1415 Pro	Ala	Gly Val	Pro	Leu 1435	1420 Asn	Pro) Phe	Phe Ile	Pro	Pro 1440
Glu 1425	1410 Tyr	Val) Val	Cys Ser	Gly	Thr 1430 Gly	1415 Pro	Ala	Gly Val	Pro	Leu 1435 Ile	1420 Asn	Pro) Phe	Phe Ile	Pro Ala	Pro 1440 Ser
Glu 1425 Gly	1410 Tyr S	Val Val Ser	Cys Ser Gln	Gly His	Thr 1430 Gly	1415 Pro	Ala His	Gly Val Thr	Pro Gly 1450	Leu 1435 Ile	1420 Asn	Pro Phe Cys	Phe Ile Gly	Pro Ala Lys	Pro 1440 Ser
Glu 1425 Gly	1410 Tyr S	Val Val Ser	Cys Ser Gln	Gly His 1445 Val	Thr 1430 Gly	1415 Pro	Ala His	Gly Val Thr	Pro Gly 1450 Gly	Leu 1435 Ile	1420 Asn	Pro Phe Cys	Phe Ile Gly	Pro Ala Lys 1455	Pro 1440 Ser
Glu 1425 Gly Met	1410 Tyr Gly Met	Val Val Ser	Cys Ser Gln Ser	Gly His 1445 Val	Thr 1430 Gly Ser	Pro Pro	Ala His Phe	Gly Val Thr Gly	Pro Gly 1450 Gly	Leu 1435 Ile Arg	1420 Asn Ala Gly	Pro Phe Cys	Phe Ile Gly Val	Pro Ala Lys 1455	Pro 1440 Ser Leu
Glu 1425 Gly Met	1410 Tyr Gly Met	Val Val Ser Arg	Cys Ser Gln Ser 1460 Asn	Gly His 1445 Val	Thr 1430 Gly Ser Val	1415 Pro Pro Leu	Ala His Phe Met Gly	Gly Val Thr Gly 1465	Pro Gly 1450 Gly Ser	Leu 1435 Ile Arg	1420 Asn Ala Gly	Pro Phe Cys Gly Ser	Phe Ile Gly Val 1470 Ser	Pro Ala Lys 1455 Pro	Pro 1440 Ser Leu Ser
Glu 1425 Gly Met	1410 Tyr Gly Met	Val Val Ser Arg 1475	Cys Ser Gln Ser 1460 Asn	Gly His 1445 Val	Thr 1430 Gly Ser Val	1415 Pro Pro Leu	Ala His Phe Met Gly 1480 Pro	Gly Val Thr Gly 1465	Pro Gly 1450 Gly Ser	Leu 1435 Ile Arg	1420 Asn Ala Gly	Pro Phe Cys Gly Ser	Phe Ile Gly Val 1470 Ser	Pro Ala Lys 1455 Pro	Pro 1440 Ser Leu Ser

Ala Thr Asp Pro Ser Leu Tyr Asn Met Asp Met Phe Tyr Ser Ser Asn Ile Pro Ala Thr Ala Arg Pro Tyr Arg Pro Tyr Ile Ile Arg Gly Met Ala Pro Pro Thr Thr Pro Cys Ser Thr Asp Val Cys Asp Ser Asp Tyr Ser Ala Ser Arg Trp Lys Ala Ser Lys Tyr Tyr Leu Asp Leu Asn Ser Asp Ser Asp Pro Tyr Pro Pro Pro Pro Thr Pro His Ser Gln Tyr Leu Ser Ala Glu Asp Ser Cys Pro Pro Ser Pro Ala Thr Glu Arg Ser Tyr Phe His Leu Phe Pro Pro Pro Pro Ser Pro Cys Thr Asp Ser Ser

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Asp	Val	Ser	Glu	Glu	Ala	Ile	Lys	Gln	Thr	Tyr	Leu	Asn	Gln	Thr	Gly
				85					90					95	
Ala	Ala	Val	Gln	Asn	Val	Val	Ile	Ser	Gly	Leu	Val	Ser	Pro	Asp	Gly
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Leu	Ala	Cys	Asp	Trp	Val	Gly	Lys	Lys	Leu	Tyr	Trp	Thr	Asp	Ser	Glu
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Thr	Asn	Arg	Ile	Glu	Val	Ala	Asn	Leu	Asn	Gly	Thr	Ser	Arg	Lys	Val
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Ala	His	Gly	Tyr	Met	Tyr	Trp	Thr	Asp	Trp	Val	Glu	Thr	Pro	Arg	Ile
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Glu	Arg	Ala	Gly	Met	Asp	Gly	Ser	Thr	Arg	Lys	Ile	Ile	Val	Asp	Ser
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Asp	Ile	Tyr	Trp	Pro	Asn	Gly	Leu	Thr	Ile	Asp	Leu	Glu	Glu	Gln	Lys
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Gln	Glu	Arg	Gln	Pro	Phe	Phe	His	Thr	Arg	Cys	Glu	Glu	Asp	Asn	Gly
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Gly	Trp	Ser	His	Leu	Cys	Leu	Leu	Ser	Pro	Ser	Glu	Pro	Phe	Tyr	Thr
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Cys	Ala	Cys	Pro	Thr	Gly	Val	Gln	Met	Gln	Asp	Asn	Gly	Arg	Thr	Cys
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Lys	Ala	Gly	Ala	Glu	Glu	Val	Leu	Leu	Leu	Ala	Arg	Arg	Thr	Asp	Leu
			340					345					350		
Arg	Arg	Ile	Ser	Leu	Asp	Thr	Pro	Asp	Phe	Thr	Asp	Ile	Val	Leu	Gln
		355					360					365			
Val	Asp	Asp	Ile	Arg	His	Ala	Ile	Ala	Ile	Asp	Tyr	Asp	Pro	Leu	Glu
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Tyr	Leu	Asp	Gly	Ser	Gly	Ala	Gln	Thr	Leu	Val	Asn	Thr	Glu	Ile	Asn
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Asp	Pro	Asp	Gly	Ile	Ala	Val	Asp	Trp	Val	Ala	Arg	Asn	Leu	Tyr	Trp
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Thr	Asp	Thr	Gly	Thr	Asp	Arg	Ile	Glu	Val	Thr	Arg	Leu	Asn	Gly	Thr
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Ser	Arg	Lys	Ile	Leu	Val	Ser	Glu	Asp	Leu	Asp	Glu	Pro	Arg	Ala	Ile
	450					455					460				
Ala	Leu	His	Pro	Val	Met	Gly	Leu	Met	Tyr	Trp	Thr	Asp	Trp	Gly	Glu
465					470					475					480
Asn	Pro	Lys	Ile	Glu	Cys	Ala	Asn	Leu	Asp	Gly	Gln	Glu	Arg	Arg	Val
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Leu	Val	Asn	Ala	Ser	Leu	Gly	Trp	Pro	Asn	Gly	Leu	Ala	Leu	Asp	Leu
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Gln	Glu	Gly	Lvs	Leu	Tvr	Trp	Glv	Asp	Ala	Lvs	Thr	Asp	Lvs	Tle	Glu

		515					520					525			
Val	Ile	Asn	Val	Asp	Gly	Thr	Lys	Arg	Arg	Thr	Leu	Leu	Glu	Asp	Lys
	530					535					540				
Leu	Pro	His	Ile	Phe	Gly	Phe	Thr	Leu	Leu	Gly	Asp	Phe	Ile	Tyr	Trp
545					550					555					560
Thr	Asp	Trp	Gln	Arg	Arg	Ser	Ile	Glu	Arg	Val	His	Lys	Val	Lys	Ala
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Gln Val Leu Val Trp Arg Asp Leu Asp Asn Pro Arg Ser Leu Ala Leu Asp Pro Thr Lys Gly Tyr Ile Tyr Trp Thr Glu Trp Gly Gly Lys Pro Arg Ile Val Arg Ala Phe Met Asp Gly Thr Asn Cys Met Thr Leu Val Asp Lys Val Gly Arg Ala Asn Asp Leu Thr Ile Asp Tyr Ala Asp Gln Arg Leu Tyr Trp Thr Asp Leu Asp Thr Asn Met Ile Glu Ser Ser Asn Met Leu Gly Gln Glu Arg Val Val Ile Ala Asp Asp Leu Pro His Pro Phe Gly Leu Thr Gln Tyr Ser Asp Tyr Ile Tyr Trp Thr Asp Trp Asn Leu His Ser Ile Glu Arg Ala Asp Lys Thr Ser Gly Arg Asn Arg Thr Leu Ile Gln Gly His Leu Asp Phe Val Met Asp Ile Leu Val Phe His Ser Ser Arg Gln Asp Gly Leu Asn Asp Cys Met His Asn Asn Gly Gln Cys Gly Gln Leu Cys Leu Ala Ile Pro Gly Gly His Arg Cys Gly Cys Ala Ser His Tyr Thr Leu Asp Pro Ser Ser Arg Asn Cys Ser Pro Pro Thr Thr Phe Leu Leu Phe Ser Gln Lys Ser Ala Ile Ser Arg Met Ile Pro Asp Asp Gln His Ser Pro Asp Leu Ile Leu Pro Leu His Gly Leu Arg Asn Val Lys Ala Ile Asp Tyr Asp Pro Leu Asp Lys Phe Ile Tyr

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<213> Homo sapiens

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<220>

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3180

4860

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<223> Identity of nucleotide sequences at the above locations are unknown.

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